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## SEQUENCE LISTING

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<120> Methods for enhancing antibody activity

<130> 14875-164US1

<150> PCT/JP2004/018493

<151> 2004-12-10

<150> JP 2003-415760

<151> 2003-12-12

<160> 28

<170> PatentIn version 3.1

<210> 1

<211> 1924

<212> DNA

<213> Macaca fascicularis

<220>

<221> CDS

<222> (11)..(1918)

<223>

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Leu Leu Ala Pro Gln Asn Leu Ala Gln Val Ser Ser Gln Asp Val Ser  
15 20 25

ttg ctg gcc tcg gac tca gag ccc ctg aag tgt ttc tcc cga aca ttt 145  
Leu Leu Ala Ser Asp Ser Glu Pro Leu Lys Cys Phe Ser Arg Thr Phe  
30 35 40 45

gag gac ctc act tgc ttc tgg gat gag gaa gag gca gca ccc agt ggg 193  
Glu Asp Leu Thr Cys Phe Trp Asp Glu Glu Glu Ala Ala Pro Ser Gly  
50 55 60

aca tac cag ctg ctg tat gcc tac ccg ggg gag aag ccc cgt gcc tgc 241  
Thr Tyr Gln Leu Leu Tyr Ala Tyr Pro Gly Glu Lys Pro Arg Ala Cys  
65 70 75

ccc ctg agt tct cag agc gtg ccc cgc ttt gga acc cga tac gtg tgc 289  
Pro Leu Ser Ser Gln Ser Val Pro Arg Phe Gly Thr Arg Tyr Val Cys  
80 85 90

cag ttt cca gcc cag gaa gaa gtg cgt ctc ttc tct ccg ctg cac ctc 337

Gln	Phe	Pro	Ala	Gln	Glu	Glu	Val	Arg	Leu	Phe	Ser	Pro	Leu	His	Leu	
95				100					105							
tgg	gtg	aag	aat	gtg	ttc	cta	aac	cag	act	cag	att	cag	cga	gtc	ctc	385
Trp	Val	Lys	Asn	Val	Phe	Leu	Asn	Gln	Thr	Gln	Ile	Gln	Arg	Val	Leu	
110				115					120						125	
ttt	gtg	gac	agt	gta	ggc	ctg	ccg	gct	ccc	ccc	agt	atc	atc	aag	gcc	433
Phe	Val	Asp	Ser	Val	Gly	Leu	Pro	Ala	Pro	Pro	Ser	Ile	Ile	Lys	Ala	
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atg	ggt	ggg	agc	cag	cca	ggg	gaa	ctt	cag	atc	agc	tgg	gag	gcc	cca	481
Met	Gly	Gly	Ser	Gln	Pro	Gly	Glu	Leu	Gln	Ile	Ser	Trp	Glu	Ala	Pro	
				145					150						155	
gct	cca	gaa	atc	agt	gat	ttc	ctg	agg	tac	gaa	ctc	cgc	tat	gcc	ccc	529
Ala	Pro	Glu	Ile	Ser	Asp	Phe	Leu	Arg	Tyr	Glu	Leu	Arg	Tyr	Gly	Pro	
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aaa	gat	ctc	aag	aac	tcc	act	ggt	ccc	acg	gtc	ata	cag	ttg	atc	gcc	577
Lys	Asp	Leu	Lys	Asn	Ser	Thr	Gly	Pro	Thr	Val	Ile	Gln	Leu	Ile	Ala	
				175					180						185	
aca	gaa	acc	tgc	tgc	cct	gct	ctg	cag	agg	cca	cac	tca	gcc	tct	gct	625
Thr	Glu	Thr	Cys	Cys	Pro	Ala	Leu	Gln	Arg	Pro	His	Ser	Ala	Ser	Ala	
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ctg	gac	cag	tct	cca	tgt	gct	cag	ccc	aca	atg	ccc	tgg	caa	gat	gga	673
Leu	Asp	Gln	Ser	Pro	Cys	Ala	Gln	Pro	Thr	Met	Pro	Trp	Gln	Asp	Gly	
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cca	aag	cag	acc	tcc	cca	act	aga	gaa	gct	tca	gct	ctg	aca	gca	gtg	721
Pro	Lys	Gln	Thr	Ser	Pro	Thr	Arg	Glu	Ala	Ser	Ala	Leu	Thr	Ala	Val	
				225					230						235	
ggt	gga	agc	tgc	ctc	atc	tca	gga	ctc	cag	cct	ggc	aac	tcc	tac	tgg	769
Gly	Gly	Ser	Cys	Leu	Ile	Ser	Gly	Leu	Gln	Pro	Gly	Asn	Ser	Tyr	Trp	
				240					245						250	
ctg	cag	ctg	cgc	agc	gaa	cct	gat	ggg	atc	tcc	ctc	ggt	ggc	tcc	tgg	817
Leu	Gln	Leu	Arg	Ser	Glu	Pro	Asp	Gly	Ile	Ser	Leu	Gly	Gly	Ser	Trp	
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gga	tcc	tgg	tcc	ctc	cct	gtg	act	gtg	gac	ctg	cct	gga	gat	gca	gtg	865
Gly	Ser	Trp	Ser	Leu	Pro	Val	Thr	Val	Asp	Leu	Pro	Gly	Asp	Ala	Val	
				270					275						280	285
gca	att	gga	ctg	caa	tgc	ttt	acc	ttg	gac	ctg	aag	aat	gtt	acc	tgt	913
Ala	Ile	Gly	Leu	Gln	Cys	Phe	Thr	Leu	Asp	Leu	Lys	Asn	Val	Thr	Cys	
				290					295						300	
caa	tgg	cag	caa	gag	gac	cat	gct	agt	tcc	caa	ggt	ttc	ttc	tac	cac	961
Gln	Trp	Gln	Gln	Glu	Asp	His	Ala	Ser	Ser	Gln	Gly	Phe	Phe	Tyr	His	
				305					310						315	
agc	agg	gca	cgg	tgc	tgc	ccc	aga	gac	agg	tac	ccc	atc	tgg	gag	gac	1009
Ser	Arg	Ala	Arg	Cys	Cys	Pro	Arg	Asp	Arg	Tyr	Pro	Ile	Trp	Glu	Asp	

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tct cgc tgc cac ttc aag tca cga aat gac agc gtt att cac atc ctt Ser Arg Cys His Phe Lys Ser Arg Asn Asp Ser Val Ile His Ile Leu 350	355	360	1105
gtg gag gtg acc aca gcc ctg ggt gct gtt cac agt tac ctg ggc tcc Val Glu Val Thr Thr Ala Leu Gly Ala Val His Ser Tyr Leu Gly Ser 370	375	380	1153
cct ttc tgg atc cac cag gct gtg cgc ctc ccc acc cca aac ttg cac Pro Phe Trp Ile His Gln Ala Val Arg Leu Pro Thr Pro Asn Leu His 385	390	395	1201
tgg agg gag atc tcc agc ggg cat ctg gaa ttg gag tgg cag cac cca Trp Arg Glu Ile Ser Ser Gly His Leu Glu Leu Glu Trp Gln His Pro 400	405	410	1249
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gaa ggc cat cag gac tgg aag gtg ctg gag ccg cct ctc ggg gcc cga Glu Gly His Gln Asp Trp Lys Val Leu Glu Pro Pro Leu Gly Ala Arg 430	435	440	1345
gga ggg acc ctg gag ctg cgc ccg cga tct cgc tac cgt tta cag ctg Gly Gly Thr Leu Glu Leu Arg Pro Arg Ser Arg Tyr Arg Leu Gln Leu 450	455	460	1393
cgc gcc agg ctc aat ggc ccc acc tac caa ggt ccc tgg agc tcg tgg Arg Ala Arg Leu Asn Gly Pro Thr Tyr Gln Gly Pro Trp Ser Ser Trp 465	470	475	1441
tcg gac cca gct agg gtg gag acc gcc acc gag acc gcc tgg att tcc Ser Asp Pro Ala Arg Val Glu Thr Ala Thr Glu Thr Ala Trp Ile Ser 480	485	490	1489
ttg gtg acc gct ctg ctg cta gtg ctg ggc ctc agc gcc gtc ctg ggc Leu Val Thr Ala Leu Leu Leu Val Leu Gly Leu Ser Ala Val Leu Gly 495	500	505	1537
ctg ctg ctg agg tgg cag ttt cct gca cac tac agg aga ctg agg Leu Leu Leu Arg Trp Gln Phe Pro Ala His Tyr Arg Arg Leu Arg 510	515	520	1585
cat gcc ctg tgg ccc tca ctt cca gat ctg cac cga gtc cta ggc cag His Ala Leu Trp Pro Ser Leu Pro Asp Leu His Arg Val Leu Gly Gln 530	535	540	1633
tac ctt agg gac act gca gcc ctg agt ccg ccc aag gcc aca gtc tca Tyr Leu Arg Asp Thr Ala Ala Leu Ser Pro Pro Lys Ala Thr Val Ser 545	550	555	1681

gat acc tgt gaa gaa gtg gaa ccc agc ctc ctt gaa atc ctc ccc aag		1729	
Asp Thr Cys Glu Glu Val Glu Pro Ser Leu Leu Glu Ile Leu Pro Lys			
560	565	570	
tcc tca gag agg act cct ttg ccc ctg tgt tcc tcc cag tcc cag atg		1777	
Ser Ser Glu Arg Thr Pro Leu Pro Leu Cys Ser Ser Gln Ser Gln Met			
575	580	585	
gac tac cga aga ttg cag cct tct tgc ctg ggg acc atg ccc ctg tct		1825	
Asp Tyr Arg Arg Leu Gln Pro Ser Cys Leu Gly Thr Met Pro Leu Ser			
590	595	600	605
gtg tgc cca ccc atg gct gag tca ggg tcc tgc tgt acc acc cac att		1873	
Val Cys Pro Pro Met Ala Glu Ser Gly Ser Cys Cys Thr Thr His Ile			
610	615	620	
gcc aac cat tcc tac cta cca cta agc tat tgg cag cag cct tga		1918	
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Ser Asp Ser Glu Pro Leu Lys Cys Phe Ser Arg Thr Phe Glu Asp Leu			
35	40	45	
Thr Cys Phe Trp Asp Glu Glu Ala Ala Pro Ser Gly Thr Tyr Gln			
50	55	60	
Leu Leu Tyr Ala Tyr Pro Gly Glu Lys Pro Arg Ala Cys Pro Leu Ser			
65	70	75	80
Ser Gln Ser Val Pro Arg Phe Gly Thr Arg Tyr Val Cys Gln Phe Pro			
85	90	95	
Ala Gln Glu Glu Val Arg Leu Phe Ser Pro Leu His Leu Trp Val Lys			
100	105	110	
Asn Val Phe Leu Asn Gln Thr Gln Ile Gln Arg Val Leu Phe Val Asp			
115	120	125	
Ser Val Gly Leu Pro Ala Pro Pro Ser Ile Ile Lys Ala Met Gly Gly			
130	135	140	
Ser Gln Pro Gly Glu Leu Gln Ile Ser Trp Glu Ala Pro Ala Pro Glu			

145	150	155	160
Ile Ser Asp Phe Leu Arg Tyr Glu Leu Arg Tyr Gly Pro Lys Asp Leu			
165	170	175	
Lys Asn Ser Thr Gly Pro Thr Val Ile Gln Leu Ile Ala Thr Glu Thr			
180	185	190	
Cys Cys Pro Ala Leu Gln Arg Pro His Ser Ala Ser Ala Leu Asp Gln			
195	200	205	
Ser Pro Cys Ala Gln Pro Thr Met Pro Trp Gln Asp Gly Pro Lys Gln			
210	215	220	
Thr Ser Pro Thr Arg Glu Ala Ser Ala Leu Thr Ala Val Gly Gly Ser			
225	230	235	240
Cys Leu Ile Ser Gly Leu Gln Pro Gly Asn Ser Tyr Trp Leu Gln Leu			
245	250	255	
Arg Ser Glu Pro Asp Gly Ile Ser Leu Gly Gly Ser Trp Gly Ser Trp			
260	265	270	
Ser Leu Pro Val Thr Val Asp Leu Pro Gly Asp Ala Val Ala Ile Gly			
275	280	285	
Leu Gln Cys Phe Thr Leu Asp Leu Lys Asn Val Thr Cys Gln Trp Gln			
290	295	300	
Gln Glu Asp His Ala Ser Ser Gln Gly Phe Phe Tyr His Ser Arg Ala			
305	310	315	320
Arg Cys Cys Pro Arg Asp Arg Tyr Pro Ile Trp Glu Asp Cys Glu Glu			
325	330	335	
Glu Glu Lys Thr Asn Pro Gly Leu Gln Thr Pro Gln Phe Ser Arg Cys			
340	345	350	
His Phe Lys Ser Arg Asn Asp Ser Val Ile His Ile Leu Val Glu Val			
355	360	365	
Thr Thr Ala Leu Gly Ala Val His Ser Tyr Leu Gly Ser Pro Phe Trp			
370	375	380	
Ile His Gln Ala Val Arg Leu Pro Thr Pro Asn Leu His Trp Arg Glu			
385	390	395	400
Ile Ser Ser Gly His Leu Glu Leu Glu Trp Gln His Pro Ser Ser Trp			
405	410	415	
Ala Ala Gln Glu Thr Cys Tyr Gln Leu Arg Tyr Thr Gly Glu Gly His			
420	425	430	
Gln Asp Trp Lys Val Leu Glu Pro Pro Leu Gly Ala Arg Gly Gly Thr			
435	440	445	
Leu Glu Leu Arg Pro Arg Ser Arg Tyr Arg Leu Gln Leu Arg Ala Arg			

450	455	460
Leu Asn Gly Pro Thr Tyr Gln Gly Pro Trp Ser Ser Trp Ser Asp Pro		
465	470	475
Ala Arg Val Glu Thr Ala Thr Glu Thr Ala Trp Ile Ser Leu Val Thr		
485	490	495
Ala Leu Leu Leu Val Leu Gly Leu Ser Ala Val Leu Gly Leu Leu Leu		
500	505	510
Leu Arg Trp Gln Phe Pro Ala His Tyr Arg Arg Leu Arg His Ala Leu		
515	520	525
Trp Pro Ser Leu Pro Asp Leu His Arg Val Leu Gly Gln Tyr Leu Arg		
530	535	540
Asp Thr Ala Ala Leu Ser Pro Pro Lys Ala Thr Val Ser Asp Thr Cys		
545	550	555
Glu Glu Val Glu Pro Ser Leu Leu Glu Ile Leu Pro Lys Ser Ser Glu		
565	570	575
Arg Thr Pro Leu Pro Leu Cys Ser Ser Gln Ser Gln Met Asp Tyr Arg		
580	585	590
Arg Leu Gln Pro Ser Cys Leu Gly Thr Met Pro Leu Ser Val Cys Pro		
595	600	605
Pro Met Ala Glu Ser Gly Ser Cys Cys Thr Thr His Ile Ala Asn His		
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Ser Tyr Leu Pro Leu Ser Tyr Trp Gln Gln Pro		
625	630	635

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<220>  
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23

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<220>  
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<222> (1)..(411)  
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1 5 10 15

gtc cac tcc cag gtt cag ctg cag cag tct gga cct gag ctg gtg aag 96  
Val His Ser Gln Val Gln Leu Gln Gln Ser Gly Pro Glu Leu Val Lys  
20 25 30

cct ggg gcc tca gtg aag att tcc tgc aag gct tct ggc tat gca ttc 144  
Pro Gly Ala Ser Val Lys Ile Ser Cys Lys Ala Ser Gly Tyr Ala Phe  
35 40 45

act aac tcc tgg atg aac tgg gtg aag cag agg cct gga aag ggt ctt 192  
Thr Asn Ser Trp Met Asn Trp Val Lys Gln Arg Pro Gly Lys Gly Leu  
50 55 60

gag tgg att gga cg att tat cct gga gat gga gaa act atc tac aat 240  
Glu Trp Ile Gly Arg Ile Tyr Pro Gly Asp Gly Glu Thr Ile Tyr Asn  
65 70 75 80

ggg aaa ttc agg gtc aag gcc aca ctg act gca gac aaa tcc tcc agc 288  
Gly Lys Phe Arg Val Lys Ala Thr Leu Thr Ala Asp Lys Ser Ser Ser  
85 90 95

aca gcc tac atg gat atc agc agc ctg aca tct gag gac tct gcg gtc 336  
Thr Ala Tyr Met Asp Ile Ser Ser Leu Thr Ser Glu Asp Ser Ala Val  
100 105 110

tac ttc tgt gca aga ggc tat gat gat tac tcg ttt gct tac tgg ggc 384  
Tyr Phe Cys Ala Arg Gly Tyr Asp Asp Tyr Ser Phe Ala Tyr Trp Gly  
115 120 125

caa ggg act ctg gtc act gtc tct gca 411  
Gln Gly Thr Leu Val Thr Val Ser Ala  
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25

30

Pro Gly Ala Ser Val Lys Ile Ser Cys Lys Ala Ser Gly Tyr Ala Phe  
 35 40 45

Thr Asn Ser Trp Met Asn Trp Val Lys Gln Arg Pro Gly Lys Gly Leu  
 50 55 60

Glu Trp Ile Gly Arg Ile Tyr Pro Gly Asp Gly Glu Thr Ile Tyr Asn  
 65 70 75 80

Gly Lys Phe Arg Val Lys Ala Thr Leu Thr Ala Asp Lys Ser Ser Ser  
 85 90 95

Thr Ala Tyr Met Asp Ile Ser Ser Leu Thr Ser Glu Asp Ser Ala Val  
 100 105 110

Tyr Phe Cys Ala Arg Gly Tyr Asp Asp Tyr Ser Phe Ala Tyr Trp Gly  
 115 120 125

Gln Gly Thr Leu Val Thr Val Ser Ala  
 130 135

&lt;210&gt; 7

&lt;211&gt; 396

&lt;212&gt; DNA

&lt;213&gt; Mus musculus

&lt;220&gt;

&lt;221&gt; CDS

&lt;222&gt; (1)..(396)

&lt;223&gt;

&lt;400&gt; 7

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 1 5 10 15

gga gcc att ggg gat att gtg atg act cag gct gca ccc tct ata cct 96  
 Gly Ala Ile Gly Asp Ile Val Met Thr Gln Ala Ala Pro Ser Ile Pro  
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gtc act cct gga gag tca gta tcc atc tcc tgt agg tct agt aag agt 144  
 Val Thr Pro Gly Glu Ser Val Ser Ile Ser Cys Arg Ser Ser Lys Ser  
 35 40 45

ctc ctg cat agt aat ggc aac act tac ttg tat tgg ttc ctg cag agg 192  
 Leu Leu His Ser Asn Gly Asn Thr Tyr Leu Tyr Trp Phe Leu Gln Arg  
 50 55 60

cca ggc cag tct cct caa ctc ctg ata tat cgg atg tcc aac ctt gcc 240  
 Pro Gly Gln Ser Pro Gln Leu Leu Ile Tyr Arg Met Ser Asn Leu Ala  
 65 70 75 80

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cgatatattgt gatgactcag gc 82

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<220>  
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<210> 18  
<211> 81  
<212> DNA  
<213> Artificial

<220>  
<223> an artificially synthesized primer sequence

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cttttatttc cagttggtc c 81

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<212> PRT  
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<400> 19

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Ser Val Lys Ile Ser Cys Arg Ala Phe Gly Tyr Ala Phe Ser Asn Ser  
 20 25 30

Trp Met Asn Trp Val Lys Gln Arg Pro Gly Lys Gly Leu Glu Trp Ile  
 35 40 45

Gly Arg Ile Tyr Pro Gly Asp Gly Glu Thr Asn Asn Asn Gly Lys Phe  
 50 55 60

Lys Gly Lys Ala Thr Leu Thr Ala Asp Lys Ser Ser Ser Thr Ala Tyr  
 65 70 75 80

Met Gln Leu Ser Ser Leu Thr Ser Glu Asp Ser Ala Val Tyr Phe Cys  
 85 90 95

Ala Arg Gly Tyr Gly Asp Tyr Ser Phe Ala Tyr Trp Gly Gln Gly Thr  
 100 105 110

Leu Val Thr Val Ser Ala  
 115

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<212> PRT

<213> Mus musculus

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 1 5 10 15

Ser Val Lys Ile Ser Cys Lys Ala Ser Gly Tyr Ala Phe Ser Ser Ser  
 20 25 30

Trp Met Asn Trp Val Lys Gln Arg Pro Gly Lys Gly Leu Glu Trp Ile  
 35 40 45

Gly Arg Ile Tyr Pro Gly Asp Gly Glu Thr Asn Asn Asn Gly Lys Phe  
 50 55 60

Lys Gly Lys Ala Thr Leu Thr Ala Asp Lys Ser Ser Thr Thr Ala Tyr  
 65 70 75 80

Met Gln Leu Ser Ser Leu Thr Ser Glu Asp Ser Ala Val Tyr Phe Cys  
 85 90 95

Ala Arg Gly Tyr Gly Asp Tyr Ser Phe Ala Tyr Trp Gly Gln Gly Thr  
 100 105 110

Leu Val Thr Val Ser Ala  
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<210> 21

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 Trp Met Asn Trp Val Lys Gln Arg Pro Gly Lys Gly Leu Glu Trp Ile  
 35 40 45  
 Gly Arg Ile Tyr Pro Gly Asp Gly Glu Thr Ile Tyr Asn Gly Lys Phe  
 50 55 60  
 Arg Val Lys Ala Thr Leu Thr Ala Asp Lys Ser Ser Ser Thr Ala Tyr  
 65 70 75 80  
 Met Asp Ile Ser Ser Leu Thr Ser Glu Asp Ser Ala Val Tyr Phe Cys  
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 35 40 45  
 Gly Arg Ile His Pro Tyr Asp Ser Glu Thr His Tyr Asn Gln Lys Phe  
 50 55 60  
 Lys Asn Lys Ala Thr Leu Thr Val Asp Lys Ser Ser Ser Thr Ala Tyr  
 65 70 75 80  
 Ile Gln Leu Ser Ser Leu Thr Ser Glu Asp Ser Ala Val Tyr Tyr Cys  
 85 90 95  
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Tyr Ala Trp Ser Trp Ile Arg Gln Leu Pro Gly Asn Lys Leu Glu Trp  
 35 40 45

Met Gly Tyr Ile Thr Tyr Ser Gly Tyr Ser Ile Tyr Asn Pro Ser Leu  
 50 55 60

Lys Ser Arg Ile Ser Ile Ser Arg Asp Thr Ser Lys Asn Gln Leu Phe  
 65 70 75 80

Leu Gln Leu Asn Ser Val Thr Thr Glu Asp Thr Ala Thr Tyr Tyr Cys  
 85 90 95

Val Gly Gly Tyr Asp Asn Met Asp Tyr Trp Gly Gln Gly Thr Ser Val  
 100 105 110

Thr Val Ser Ser  
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 1 5 10 15

Glu Ser Val Ser Ile Ser Cys Arg Ser Ser Lys Ser Leu Leu His Ser  
 20 25 30

Asn Gly Asn Thr Tyr Leu Tyr Trp Phe Leu Gln Arg Pro Gly Gln Ser  
 35 40 45

Pro Gln Leu Leu Ile Tyr Arg Met Ser Asn Leu Ala Ser Gly Val Pro  
 50 55 60

Asp Arg Phe Ser Gly Ser Gly Ser Gly Ala Ala Phe Thr Leu Arg Ile  
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Ser Arg Val Glu Ala Glu Asp Val Gly Val Tyr Tyr Cys Met Gln His  
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Leu Glu Tyr Pro Tyr Thr Phe Gly Ser Gly Thr Lys Leu Glu Ile Lys  
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35 40 45  
Pro Gln Leu Leu Ile Tyr Arg Met Ser Asn Leu Ala Ser Gly Val Pro  
50 55 60  
Asp Arg Phe Ser Gly Ser Gly Ala Ala Phe Thr Leu Arg Ile  
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20 25 30  
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35 40 45  
Pro Gln Leu Leu Ile Tyr Arg Met Ser Asn Leu Ala Ser Gly Val Pro  
50 55 60  
Asp Arg Phe Ser Gly Ser Gly Thr Ala Phe Thr Leu Arg Ile  
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Asp	Ile	Val	Met	Thr	Gln	Ala	Ala	Pro	Ser	Val	Pro	Val	Thr	Pro	Gly
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Glu	Ser	Val	Ser	Ile	Ser	Cys	Arg	Ser	Ser	Lys	Ser	Leu	Leu	Tyr	Ser
					20				25				30		

Asn	Gly	Asn	Thr	Tyr	Leu	Tyr	Trp	Phe	Leu	Gln	Arg	Pro	Gly	Gln	Ser
							35		40			45			

Pro	Gln	Leu	Leu	Ile	Tyr	Arg	Met	Ser	Asn	Leu	Ala	Ser	Gly	Val	Pro
						50		55			60				

Asp	Arg	Phe	Ser	Gly	Ser	Gly	Ser	Gly	Thr	Ala	Phe	Thr	Leu	Thr	Ile
					65		70		75			80			

Ser	Ser	Val	Glu	Ala	Glu	Asp	Val	Gly	Val	Tyr	Tyr	Cys	Met	Gln	His
					85			90				95			

Leu	Glu	Tyr	Pro	Tyr	Thr	Phe	Gly	Ser	Gly	Thr	Lys	Leu	Glu	Ile	Lys
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<213> Mus musculus

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Glu	Lys	Val	Thr	Leu	Thr	Cys	Ser	Ala	Ser	Ser	Ser	Val	Ser	Ser	Ser
					20			25			30				

His	Leu	Tyr	Trp	Tyr	Gln	Gln	Lys	Pro	Gly	Ser	Ser	Pro	Lys	Leu	Trp
					35		40			45					

Ile	Tyr	Ser	Thr	Ser	Asn	Leu	Ala	Ser	Gly	Val	Pro	Ala	Arg	Phe	Ser
					50		55			60					

Gly	Ser	Gly	Ser	Gly	Thr	Ser	Tyr	Ser	Leu	Thr	Ile	Ser	Asn	Met	Glu
					65		70		75		80				

Thr	Glu	Asp	Ala	Ala	Ser	Tyr	Phe	Cys	His	Gln	Trp	Ser	Ser	Tyr	Pro
					85			90			95				

Trp	Thr	Phe	Gly	Gly	Thr	Lys	Leu	Glu	Ile	Lys				
					100		105							